
Sequence Listing was accepted.

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Reviewer: markspencer

Timestamp: [year=2008; month=7; day=28; hr=9; min=40; sec=6; ms=230;]

Validated By CRFValidator v 1.0.3

Application No: 10516317 Version No: 2.0

Input Set:

Output Set:

Started: 2008-07-25 21:38:45.411 **Finished:** 2008-07-25 21:38:47.803

Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 392 ms

Total Warnings: 22
Total Errors: 0

No. of SeqIDs Defined: 22
Actual SeqID Count: 22

Error code Error Description

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W	402	Undefined organism found in <213> in SEQ ID (2)	
W	213	Artificial or Unknown found in <213> in SEQ ID	(3)
W	213	Artificial or Unknown found in <213> in SEQ ID	(4)
W	213	Artificial or Unknown found in <213> in SEQ ID	(5)
W	213	Artificial or Unknown found in <213> in SEQ ID	(6)
W	213	Artificial or Unknown found in <213> in SEQ ID	(7)
W	213	Artificial or Unknown found in <213> in SEQ ID	(8)
W	213	Artificial or Unknown found in <213> in SEQ ID	(9)
W	213	Artificial or Unknown found in <213> in SEQ ID	(10)
W	402	Undefined organism found in <213> in SEQ ID (11	_)
W	402	Undefined organism found in <213> in SEQ ID (12	?)
W	402	Undefined organism found in <213> in SEQ ID (13	3)
W	402	Undefined organism found in <213> in SEQ ID (14	l)
W	402	Undefined organism found in <213> in SEQ ID (15	5)
W	402	Undefined organism found in <213> in SEQ ID (16	5)
W	402	Undefined organism found in <213> in SEQ ID (17	⁷)
W	402	Undefined organism found in <213> in SEQ ID (18	3)
W	402	Undefined organism found in <213> in SEQ ID (19))
W	402	Undefined organism found in <213> in SEQ ID (20))

Input Set:

Output Set:

Started: 2008-07-25 21:38:45.411

Finished: 2008-07-25 21:38:47.803

Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 392 ms

Total Warnings: 22

Total Errors: 0

No. of SeqIDs Defined: 22

Actual SeqID Count: 22

Error code	Error Description
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W 402 Undefined organism found in <213> in SEQ ID (21)

 \mathbb{W} 402 Undefined organism found in <213> in SEQ ID (22)

SEQUENCE LISTING

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			02-16 06-16		3										
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Gly	Thr	Val	Asn 20	Asn	His	His	Phe	Met 25	Суз	Glu	Ala	Glu	Gly 30	Glu	Gly
Lys	Pro	Tyr 35	Glu	Gly	Thr	Gln	Met 40	Glu	Asn	Ile	Lys	Val 45	Thr	Lys	Gly
Gly	Pro 50	Leu	Pro	Phe	Ser	Phe 55	Asp	Ile	Leu	Thr	Pro 60	Asn	Cys	Gln	Tyr
Gly 65	Ser	Val	Ala	Ile	Thr 70	Lys	Tyr	Thr	Ser	Gly 75	Ile	Pro	Asp	Tyr	Phe 80
Lys	Gln	Ser	Phe	Pro 85	Glu	Gly	Phe	Thr	Trp 90	Glu	Arg	Thr	Thr	Ile 95	Tyr
Glu	Asp	Gly	Ala 100	Tyr	Leu	Thr	Thr	Gln 105	Gln	Glu	Thr	Lys	Leu 110	Asp	Gly
Asn	Суз	Leu 115	Val	Tyr	Asn	Ile	Lys 120	Ile	Leu	Gly	Суз	Asn 125	Phe	Pro	Pro
Asn	Gly 130	Pro	Val	Met	Gln	Lys 135	Lys	Thr	Gln	Gly	Trp 140	Glu	Pro	Суз	Суз

145	Met	Arg	Tyr	Thr	Arg 150	Asp	Gly	Val	Leu	Cys 155	Gly	Gln	Thr	Leu	Met 160	
Ala	Leu	Lys	Суз	Ala 165	Asp	Gly	Asn	His	Leu 170	Thr	Суз	His	Leu	Arg 175	Thr	
Thr	Tyr	Arg	Ser 180	Lys	Lys	Ala	Ala	Lys 185	Ala	Leu	Gln	Met	Pro 190	Pro	Phe	
His	Phe	Ser 195	Asp	His	Arg	Pro	Glu 200	Ile	Val	Lys	Val	Ser 205	Glu	Asn	Gly	
Thr	Leu 210	Phe	Glu	Gln	His	Glu 215	Ser	Ser	Val	Ala	Arg 220	Tyr	Суз	Gln	Thr	
Cys 225	Pro	Ser	Lys	Leu	Gly 230	His	Asn									
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	0> 1> CI 2> (1		(696))												
< 400	0> 2															
atg	0> 2 gct Ala				-	-		-	-		-			_		48
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atg Met 1 ggc Gly	gct Ala aca	Ser gtc Val	Lys aac Asn 20 gag	Ile 5 aat Asn	Ser cat His	Asp cac His	Asn ttc Phe	Val atg Met 25	Arg 10 tgc Cys	Ile gaa Glu ata	Lys gct Ala	Leu gaa Glu gtc	Tyr gga Gly 30	Met 15 gag Glu	Glu ggc Gly gga	
atg Met 1 ggc Gly aag Lys	gct Ala aca Thr	gtc Val tac Tyr 35	Lys aac Asn 20 gag Glu ccg	Ile 5 aat Asn gga Gly	Ser cat His act Thr	Asp cac His caa Gln	Asn ttc Phe atg Met 40 gat	Val atg Met 25 gag Glu atc	Arg 10 tgc Cys aac Asn	gaa Glu ata Ile	Lys gct Ala aaa Lys cct	gaa Glu gtc Val 45	gga Gly 30 acc Thr	Met 15 gag Glu aaa Lys	ggc Gly gga Gly	96
atg Met 1 ggc Gly aag Lys ggc Gly	gct Ala aca Thr cca Pro	gtc Val tac Tyr 35 ctg Leu	aac Asn 20 gag Glu ccg Pro	Ile 5 aat Asn gga Gly ttc Phe	cat His act Thr tct Ser	Asp cac His caa Gln ttt Phe 55	ttc Phe atg Met 40 gat Asp	Val atg Met 25 gag Glu atc Ile	Arg 10 tgc Cys aac Asn ttg Leu	gaa Glu ata Ile acg Thr	gct Ala aaa Lys cct Pro 60 att	gaa Glu gtc Val 45 aac Asn	gga Gly 30 acc Thr	Met 15 gag Glu aaa Lys caa Gln	Glu ggc Gly gga Gly tat Tyr	96
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aat tgc ctc gtc tac aat att aaa atc ctt gga tgt aat ttt ccc ccc
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Asn Cys Leu Val Tyr Asn Ile Lys Ile Leu Gly Cys Asn Phe Pro Pro
        115
                            120
aat ggt cct gtg atg cag aag aaa acc caa ggc tgg gaa ccc tgt tgc
Asn Gly Pro Val Met Gln Lys Lys Thr Gln Gly Trp Glu Pro Cys Cys
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                        135
gag atg cgc tat aca cgt gat ggt gtg cta tgt ggc caa aca tta atg
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Glu Met Arg Tyr Thr Arg Asp Gly Val Leu Cys Gly Gln Thr Leu Met
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145
                    150
gca ctt aaa tgc gcc gat ggg aac cac ctc act tgc cat ctg aga act
                                                                   528
Ala Leu Lys Cys Ala Asp Gly Asn His Leu Thr Cys His Leu Arg Thr
act tac agg tcc aaa aag gca gca aag gcg ttg cag atg cca ccc ttc
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Thr Tyr Arg Ser Lys Lys Ala Ala Lys Ala Leu Gln Met Pro Pro Phe
            180
                                185
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His Phe Ser Asp His Arg Pro Glu Ile Val Lys Val Ser Glu Asn Gly
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                            200
aca cta ttt gaa cag cac gaa agt tca gtg gcc agg tac tgt caa aca
                                                                   672
Thr Leu Phe Glu Gln His Glu Ser Ser Val Ala Arg Tyr Cys Gln Thr
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Cys Pro Ser Lys Leu Gly His Asn
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36

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36

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Gly Thr Val Asn Asn His His Phe Met Cys Glu Ala Glu Gly Glu Gly
20 25 30

Lys Pro Tyr Glu Gly Thr Gln Met Glu Asn Ile Lys Val Thr Lys Gly \$45\$

Gly Pro Leu Pro Phe Ser Phe Asp Ile Leu Thr Pro Asn Cys Gln Leu 50 55 60

Gly Ser Val Ala Ile Thr Lys Tyr Thr Ser Gly Ile Pro Asp Tyr Phe
65 70 75 80

Lys Gln Ser Phe Pro Glu Gly Phe Thr Trp Glu Arg Thr Thr Ile Tyr

85 90 95

Glu Asp Gly Ala Tyr Leu Thr Thr Gln Gln Glu Thr Lys Leu Asp Gly 100 105 110

Asn Cys Leu Val Tyr Asn Ile Lys Ile Leu Gly Cys Asn Phe Pro Pro 115 120 125

Asn Gly Pro Val Met Gln Lys Lys Thr Gln Gly Trp Glu Pro Cys Cys 130 135 140

Ala Leu Lys Cys Ala Asp Gly Asn His Leu Thr Cys His Leu Arg Thr 165 170 175

Thr Tyr Arg Ser Lys Lys Ala Ala Lys Ala Leu Gln Met Pro Pro Phe 180 185 190

His Phe Ser Asp His Arg Pro Glu Ile Val Lys Val Ser Glu Asn Gly
195 200 205

Thr Leu Phe Glu Gln His Glu Ser Ser Val Ala Arg Tyr Cys Gln Thr 210 215 220

165

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170

175

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aca cta ttt gaa cag cac gaa agt tca gtg gcc agg tac tgt caa aca Thr Leu Phe Glu Gln His Glu Ser Ser Val Ala Arg Tyr Cys Gln Thr 210 215 220	572										
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Gly Pro Leu Pro Phe Ser Phe Asp Ile Leu Thr Pro Asn Cys Gln Met 50 55 60											
Gly Ser Val Ala Ile Thr Lys Tyr Thr Ser Gly Ile Pro Asp Tyr Phe 65 70 75 80											
Lys Gln Ser Phe Pro Glu Gly Phe Thr Trp Glu Arg Thr Thr Ile Tyr 85 90 95											
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His Phe Ser	Asp His Arg	Pro Glu Ile 200	Val Lys Val	Ser Glu Asn 205	Gly
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Cys Pro Ser 225	Lys Leu Gly 230	His Asn			
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	-	-	tca ggg att Ser Gly Ile 75	-	
-	-		tgg gaa aga Trp Glu Arg 90		
	-		caa gaa acc Gln Glu Thr	-	
-	_		ctt gga tgt Leu Gly Cys		
		_	caa ggc tgg Gln Gly Trp 140		_

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Ala	Leu	Lys	Суз	Ala	Asp	Gly	Asn	His	Leu	Thr	Суз	His	Leu	Arg	Thr	
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Thr	Tyr	Arg	Ser	Lys	Lys	Ala	Ala	Lys	Ala	Leu	Gln	M				